

Methane Emission from a Simulated Hypersaline Microbial Mat

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We present here a simulation model of methanogenesis in hypersaline mats. This represents a new addition to our current published model (MBGC-Microbial BioGeoChemistry) that simulates O₂, DIC, and sulfide cycling by cyanobacteria and sulfur bacteria (Decker *et al.* (in press) FEMS Microbiology Ecology). Our approach is to use Michaelis-Menten dynamics to describe limits imposed on methanogenesis by substrates, environmental parameters, and other bioactive chemicals. These can be described by the equation:

$$f(\text{MET}) = V_{\text{max}} * f(\text{methylated substrates, 'MS'}) * f(\text{O}_2) * f(\text{salinity}) * f(\text{temperature});$$

where v_{max} is the maximum reported methanogenesis rate given unlimited supply of methanol and methylamines as substrates (MS) that are not used by sulfate-reducing bacteria (SRB). We assume that MS have a saturating relationship to methanogenesis in MBGC, and is the sole substrate group for methanogenesis in this hypersaline mat where [SO₄⁻²] does not limit SRB activity. MS are derived from the osmoregulators glycine betaine and choline. MS are accumulated as compatible solutes by cyanobacteria, then released when the cyanobacteria die and decompose. The supply rate of MS to methanogens within the mat is currently unknown. In order to model MS supply to methanogens within each 0.5mm mat layer, we use greenhouse and literature values of

methanogenesis under near-optimal environmental and chemical conditions in hypersaline mats and work backwards. Salinity and temperature affect methanogenesis via optimum functions with distinct maximums. The methanogen function is then used in a larger equation that includes biomass and mat physical properties to model CH₄ production. Methane then diffuses up through the diffusive boundary layer and out of the mat.

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